JUL 2 7 2003

AMENDMENTS TO THE DRAWINGS:

Please amend FIG. 17A-C as shown in the enclosed marked-up copy labeled "Annotated Marked-up Drawings", wherein the changes of "CRYCBB2" to --CRY3BB2-- in two places in FIG. 17A; "CRYS3BB2" to --CRY3BB2-- in FIG. 17B; and the additions of -- (SEQ ID NO:109)--, --(SEQ ID NO:110)--, --(SEQ ID NO:111)--, --(SEQ ID NO:112)-- and --(SEQ ID NO:113)-- at the end of the sequences for CRY3C, CRY3BB2, CRY3BB, CRY3BA, and CRY3A, respectively, in FIG. 17C are shown in red. A replacement copy of FIG. 17A-C labeled "Replacement Sheet" is also provided as an appendix to this amendment.

Response to Office Action 10/614,076

replacement pages 2 of 10, 3 of 10, and 7 of 10 to substitute for pages 2 of 10, 3 of 10, and 7 of 10 of the previous Response filed May 9, 2005.

REMARKS

This paper is filed in response to the Notice of Non-Compliant Amendment. It is believed that the response is filed timely and no fees are due at this time. However, should any fees under 37 C.F.R. §§ 1.16 to 1.21 be required for any reason relating to the enclosed materials, the Commissioner is authorized to deduct said fees from Deposit Account No. 01-2508/11792.0218.DVUS01.

Respectfully submitted,

J. Wendy Davis Reg. No. 46,393

Agent for Assignee

MONSANTO TECHNOLOGY LLC

Customer No. 45607 Howrey Simon Arnold & White, LLP 2941 Fairview Park Drive, Box 7 Falls Church, VA 22042 (713) 787-1512 (direct line) (703) 336-6950 (fax)

Date:

July 25, 2005

REMARKS

Claim Status

The active claims in this case are claims 38-46. Each of claims 38 and 39 is amended

to replace "comprising" with --wherein the method comprises-- as helpfully suggested by the

Examiner. Claims 34, 36 and 37 are cancelled. As noted by the Examiner the amendments

to claim 34 in the preliminary amendment dated 7 July 2003 were not properly marked.

Applicants have therefore cancelled claim 34 (and dependant claims 36 and 37) and rewritten

them as new claims 44-46. Claims 41-43 have been amended to recite "said gene encoding

said modified Cry3Bb* polypeptide". No new matter has been introduced into the claims.

Amendments to the Drawings and Specification

FIG. 17A-C and the specification at page 58, lines 1-6 have been amended to add

sequence identifiers to the corresponding sequences that are referenced therein to fully

comply with the sequence rules. Furthermore, FIG. 17A-C has been amended to correct a

couple of typographic errors. No new matter is introduced.

Objections to Claims 34 and 38-39

Claim 34 was objected for the term "comprising". In response, new claim 44 (which

replaces claim 34) has been amended to include the phrase "wherein the method comprises"

as helpfully suggested by the Examiner. Similarly, claims 38-39 were objected for the term

"comprising". In response, claims 38-39 have been amended to include the phrase "wherein

the method comprises" as helpfully suggested by the Examiner.

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Replacement page

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ALIGNMENT OF CRY3 SEQUENCES

(Numbered according to Cry3BB) (alpha helices underlined, beta sheets marked with +++'s)

CRY3BB2	CRY3C: CRYCBB2: CRY3BB: CRY3BA: CRY3A:	1 10 20 30 40 MNPNNRSEHDTIKATENNEVSNNHAQYPLADTPTLEELNY MNPNNRSEHDTIKVTPNSELPTNHNQYPLADNPNSTLEELNY MNPNNRSEHDTIKVTPNSELQTNHNQYPLADNPNSTLEELNY MIRMGGRKMNPNNRSEYDTIKVTPNSELPTNHNQYPLADNPNSTLEELNY MIRKGGRKMNPNNRSEHDTIKTTENNEVPTNHVQYPLAETPNPTLEDLNY
CRY3BB2	CRY3C: CRYCBB2: CRY3BB: CRY3BA: CRY3A:	50 60 70 80 90 KEFLRRTTDNNVEALDSSTTKDAIQKGISIIGDLLGVVGFPYGGALVSFY KEFLRMTEDSSTEVLDNSTVKDAVGTGISVVGQILGVVGVPFAGALTSFY KEFLRMTEDSSTEVLDNSTVKDAVGTGISVVGQILGVVGVPFAGALTSFY KEFLRMTADNSTEVLDSSTVKDAVGTGISVVGQILGVVGVPFAGALTSFY KEFLRMTADNNTEALDSSTTKDVIQKGISVVGDLLGVVGFPFGGALVSFY
	CRY3C: CRY3BB2: CRY3BB: CRY3BA: CRY3A:	100 110 120 130 140 TNLLNTIWPGE-DPLKAFMQQVEALIDQKIADYAKDKATAELQGLKNVFI QSFLDTIWPSDADPWKAFMAQVEVLIDKKIEEYAKSKALAELQGLQNNFI QSFLNTIWPSDADPWKAFMAQVEVLIDKKIEEYAKSKALAELQGLQNNFI QSFLNAIWPSDADPWKAFMAQVEVLIDKKIEEYAKSKALAELQGLQNNFI TNFLNTIWPSE-DPWKAFMEQVEALMDQKIADYAKNKALAELQGLQNNVI
•	CRY3C: CRY3BB2: CRY3BB: CRY3BA: CRY3A:	150 160 170 180 190 DYVSALDSWDKTPLTLRDGRSQGRIRELFSQAESHFRRSMPSFAVSGYEV DYVNALNSWKKTPLSLRSKRSQDRIRELFSQAESHFRNSMPSFAVSKFEV DYVNALNSWKKTPLSLRSKRSQDRIRELFSQAESHFRNSMPSFAVSKFEV DYVNALDSWKKAPVNLRSRRSQDRIRELFSQAESHFRNSMPSFAVSKFEV DYVSALSSWQKNPVSSRNPHSQGRIRELFSQAESHFRNSMPSFAISGYEV

	CRY3C: CRY3BB2: CRY3BB: CRY3BA: CRY3A:	200 210 220 230 240 LFLPTYAQAANTHLLLKDAQIYGTDWGYSTDDLNEFHTKQKDLTIEYTN LFLPTYAQAANTHLLLKDAQVFGEEWGYSSEDVAEFYHRQLKLTQQYTD LFLPTYAQAANTHLLLKDAQVFGEEWGYSSEDVAEFYHRQLKLTQQYTD LFLPTYAQAANTHLLLKDAQVFGEEWGYSSEDIAEFYQRQLKLTQQYTD LFLTTYAQAANTHLFLLKDAQIYGEEWGYEKEDIAEFYKRQLKLTQEYTD)))
`CRY3BB2	CRY3C: CRYS3BB2 : CRY3BB: CRY3BA: CRY3A:	250 260 270 280 290 HCAKWYKAGLDKLRGSTYEEWVKFNRYRREMTLTVLDLITLFPLYDVRTY HCVNWYNVGLNGLRGSTYDAWVKFNRFRREMTLTVLDLIVLFPFYDVRLY HCVNWYNVGLNGLRGSTYDAWVKFNRFRREMTLTVLDLIVLFPFYDIRLY HCVNWYNVGLNSLRGSTYDAWVKFNRFRREMTLTVLDLIVLFPFYDVRLY HCVKWYNVGLDKLRGSSYESWVNFNRYRREMTLTVLDLIALFPL	, ,
	CRY3C: CRY3BB2: CRY3BB: CRY3BA: CRY3A:	300 310 320 330 340 TKGVKTELTRDVLTDPIVAVNNMNGYGTTFSNIENYIRKPHLFDYLHAIQ SKGVKTELTRDIFTDPIFSLNTLQEYGPTFLSIENSIRKPHLFDYLQGIE SKGVKTELTRDIFTDPIFSLNTLQEYGPTFLSIENSIRKPHLFDYLQGIE SKGVKTELTRDIFTDPIFTLNALQEYGPTFSSIENSIRKPHLFDYLRGIE PKEVKTELTRDVLTDPIVGVNNLRGYGTT <u>FSNIENY</u> IRKPHLFDYLHRIQ ++++++++++++++++++++++++++++++++++++	}
	CRY3C: CRY3BB2: CRY3BB: CRY3BA: CRY3A:	350 360 370 380 390 FHSRLQPGYFGTDSFNYWSGNYVSTRSSIGSDEIIRSPFYGNKSTLDVQN FHTRLQPGYSGKDSFNYWSGNYVETRPSIGSSKTITSPFYGDKSTEPVQK FHTRLQPGYFGKDSFNYWSGNYVETRPSIGSSKTITSPFYGDKSTEPVQK FHTRLRPGYSGKDSFNYWSGNYVETRPSIGSNDTITSPFYGDKSIEPIQK FHTRFQPGYYGNDSFNYWSGNYVSTRPSIGSNDIITSPFYGNKSSEPVQN ++++++++++++++++++++++++++++++++++++	
	CRY3C: CRY3BB2: CRY3BB: CRY3BA: CRY3A:	400 410 420 430 LEFNGEKVFRAVANGNLAVWPVGTGGTKIHSGVTKVQFSQYNDRKDEVRT LSFDGQKVYRTIANTDVAAWPNGKIYFGVTKVDFSQYDDQKNETST LSFDGQKVYRTIANTDVAAWPNGKIYFGVTKVDFSQYDDQKNETST LSFDGQKVYRTIANTDIAAFPDGKIYFGVTKVDFSQYDDQKNETST LEFNGEKVYRAVANTNLAVWPSAVYSGVTKVEFSQYNDQTDEAST	•

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470
                                                    480
          440
                    450
                               460
          QTYDSKRNVGGIV-FDSIDQLPPITTDESLEKAYSHQLNYVRCFLLQGGR
CRY3C:
          QTYDSKRNNGHVGAQDSIDQLPPETTDEPLEKAYSHQLNYAECFLMQDRR
CRY3BB2:
          QTYDSKRNNGHVSAQDSIDQLPPETTDEPLEKAYSHQLNYAECFLMQDRR
CRY3BB:
          OTYDSKRYNGYLGAODSIDQLPPETTDEPLEKAYSHQLNYAECFLMQDRR
CRY3BA:
          QTYDSKRNVGAVS-WDSIDQLPPETTDEPLEKGYSHQLNYVMCFLMQGSR
CRY3A:
          ++++
                       +++++
                                                  ++++++
          490
                   500
                              510
                                        520
                                                   530
          GIIPVFTWTHKSVDFYNTLDSEKITQIPFVKAFILVNSTSVVAGPGFTGG
CRY3C:
          GTIPFFTWTHRSVDFFNTIDAEKITQLPVVKAYALSSGASIIEGPGFTGG
CRY3BB2:
          GTIPFFTWTHRSVDFFNTIDAEKITOLPVVKAYALSSGASIIEGPGFTGG
CRY3BB:
CRY3BA:
          GTIPFFTWTHRSVDFFNTIDAEKITQLPVVKAYALSSGASIIEGPGFTGG
CRY3A:
          GTIPVLTWTHKSVDFFNMIDSKKITQLPLVKAYKLQSGASVVAGPRFTGG
                                            ++++++
             ++++++
                           +++++ ++++
                               560
                                         570
                                                    580
          540
                     550
          DII-KCT-NGSGLTLYVTPAPDLTYSKTYKIRIRYASTSQVRFGIDLGSY
CRY3C:
          NLLFLKESSNSIAKFKVTL-NSAALLQRYRVRIRYASTTNLRLFVQNSNN
CRY3BB2:
          NLLFLKESSNSIAKFKVTL-NSAALLQRYRVRIRYASTTNLRLFVQNSNN
CRY3BB:
          NLLFLKESSNSIAKFKVTL-NSAALLQRYRVRIRYASTTNLRLFVQNSNN
CRY3BA:
          DII-OCTENGSAATIYVTPD--VSYSQKYRARIHYASTSQITFTLSLDGA
CRY3A:
                     ++++++++
                                    +++++++++++
                                                    ++++++
                                                    630
          590
                    600
                               610
                                         620
          THSISYFDKTMDKGNTLTYNSFNLSSVSRPIEISG-GNKIGVSVGGIGSG
CRY3C:
          DFIVIYINKTMNIDDDLTYQTFDLATTNSNMGFSGDTNELIIGAESFVSN
CRY3BB2:
          DFLVIYINKTMNKDDDLTYQTFDLATTNSNMGFSGDKNELIIGAESFVSN
CRY3BB:
          DFLVIYINKTMNIDGDLTYQTFDFATSNSNMGFSGDTNDFIIGAESFVSN
CRY3BA:
          PFNQYYFDKTINKGDTLTYNSFNLASFSTPFELSG--NNLQIGVTGLSAG
CRY3A:
            ++++++
                               ++++++
          640
                    650
          DEVYIDKIEFIPMD ( SEQ ID No:109)
CRY3C:
CRY3BB2:
          EKIYIDKIEFIPVQL (SEQ ID NO:1/0)
CRY3BB:
          EKIYIDKIEFIPVQL (SEQ IDM:111)
          EKIYIDKIEFIPVQ (SEQ JO NO:1/2)
CRY3BA:
CRY3A:
          DKVYIDKIEFIPVN
                          (SEQ ID NO:113)
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